

Zagmount

1649

11

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RAW SEQUENCE LISTING
PATENT APPLICATION US/08/989,881A

DATE: 08/06/1999
TIME: 15:22:49

Input Set: H989881A.RAW

This Raw Listing contains the General Information
Section and up to first 5 pages.

ENTERED

1 <110> APPLICANT: Sheen, Jen
2 <120> TITLE OF INVENTION: STRESS-PROTECTED TRANSGENIC PLANTS
3 <130> FILE REFERENCE: 08472/716002
4 <140> CURRENT APPLICATION NUMBER: US/08/989,881A
5 <141> CURRENT FILING DATE: 1997-12-12
6 <150> EARLIER APPLICATION NUMBER: 60/032,966
7 <151> EARLIER FILING DATE: 1996-12-13
8 <160> NUMBER OF SEQ ID NOS: 18
9 <170> SOFTWARE: FastSEQ for Windows Version 3.0
10 <210> SEQ ID NO 1
11 <211> LENGTH: 1020
12 <212> TYPE: DNA
13 <213> ORGANISM: Arabidopsis thaliana
14 <220> FEATURE:
15 <221> NAME/KEY: variation
16 <222> LOCATION: (1)...(1020)
17 <223> OTHER INFORMATION: N is any nucleotide.
18 <400> SEQUENCE: 1

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20 caccgcggtg gcgccgctc tagaactagt ggatcc atg gct aat caa act cag 114
21 atc agc gac aag tac atc tta gga cga gaa ctc ggt cgc ggc gaa ttc 162
22 gga atc acg tat ctt tgt aca gat aga gag act cgt gaa gct tta gct 210
23 tgc aaa tca atc tcc aag aga aag ctc cga acc gcc gtc gat gtg gaa 258
24 gac gtc cgt cgt gaa gtc acg atc atg tca act tta ccg gaa cac cca 306
25 aac gtt gtg aaa ctt aaa gcg act tat gag gat aac gag acc gtg cat 354
26 ctt gtg atg gag ctt tgt gaa gga ggt gag ctt ttt ggt cgg att gtt 402
27 gca aga gga cat tat aca gag cgt gcg gcg gct acc gtc gcg aga acg 450
28 atc gcg gaa gtt gtg agg atg tgt cat gtc aat ggt gtt atg cat aga 498
29 gat ttg aag cct gag aat ttc ttg ttt gct aac aag aag gag aat tct 546
30 gca ctt aag gct att gat ttt ggt tta tct gtt ctc ttt aaa cct gga 594
31 gag agg ttt aca gag att gtt gga agt cct tat tat atg gct cca gaa 642
32 gtg ttg aag aga aat tat gga cca gag gtt gat gtg tgg agt gct gga 690
33 gtt atc ctc tac atc ttg ctt tgt ggt gtt cct ccg ttt tgg gca gag 738
34 act gaa caa ggt gtg gct ctt gcc atc ttg agg gga gtt ctt gat ttt 786
35 aag aga gat cct tgg tgc cag ata tca gag agc gca aag agc ctt gtg 834
36 aag cag atg ttg gaa cct gat tca act aag cgt ttg act gct cag caa 882
37 gtt ctt gat cac cct tgg ata cag aat gca aag aaa aggatcaagc 928
W--> 38 ttatcgatac cgtcgacctc gagggggggc cgggtaccag ctttngttcc ctttagtgag 988
39 ggttaatttc gagcttggcg taatcatgtc at 1020

40 <210> SEQ ID NO 2
41 <211> LENGTH: 274
42 <212> TYPE: PRT
43 <213> ORGANISM: Arabidopsis thaliana
44 <400> SEQUENCE: 2

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45      Met Ala Asn Gln Thr Gln Ile Ser Asp Lys Tyr Ile Leu Gly Arg Glu
46      1              5              10              15
47      Leu Gly Arg Gly Glu Phe Gly Ile Thr Tyr Leu Cys Thr Asp Arg Glu
48              20              25              30
49      Thr Arg Glu Ala Leu Ala Cys Lys Ser Ile Ser Lys Arg Lys Leu Arg
50              35              40              45
51      Thr Ala Val Asp Val Glu Asp Val Arg Arg Glu Val Thr Ile Met Ser
52              50              55              60
53      Thr Leu Pro Glu His Pro Asn Val Val Lys Leu Lys Ala Thr Tyr Glu
54      65              70              75              80
55      Asp Asn Glu Thr Val His Leu Val Met Glu Leu Cys Glu Gly Gly Glu
56              85              90              95
57      Leu Phe Gly Arg Ile Val Ala Arg Gly His Tyr Thr Glu Arg Ala Ala
58              100             105             110
59      Ala Thr Val Ala Arg Thr Ile Ala Glu Val Val Arg Met Cys His Val
60              115             120             125
61      Asn Gly Val Met His Arg Asp Leu Lys Pro Glu Asn Phe Leu Phe Ala
62              130             135             140
63      Asn Lys Lys Glu Asn Ser Ala Leu Lys Ala Ile Asp Phe Gly Leu Ser
64      145             150             155             160
65      Val Leu Phe Lys Pro Gly Glu Arg Phe Thr Glu Ile Val Gly Ser Pro
66              165             170             175
67      Tyr Tyr Met Ala Pro Glu Val Leu Lys Arg Asn Tyr Gly Pro Glu Val
68              180             185             190
69      Asp Val Trp Ser Ala Gly Val Ile Leu Tyr Ile Leu Leu Cys Gly Val
70              195             200             205
71      Pro Pro Phe Trp Ala Glu Thr Glu Gln Gly Val Ala Leu Ala Ile Leu
72              210             215             220
73      Arg Gly Val Leu Asp Phe Lys Arg Asp Pro Trp Ser Gln Ile Ser Glu
74      225             230             235             240
75      Ser Ala Lys Ser Leu Val Lys Gln Met Leu Glu Pro Asp Ser Thr Lys
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77      Arg Leu Thr Ala Gln Gln Val Leu Asp His Pro Trp Ile Gln Asn Ala
78              260             265             270
79      Lys Lys

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83      <213> ORGANISM: Arabidopsis thaliana
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87      <211> LENGTH: 28
88      <212> TYPE: DNA
89      <213> ORGANISM: Arabidopsis thaliana
90      <400> SEQUENCE: 4
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92      <210> SEQ ID NO 5
93      <211> LENGTH: 33
94      <212> TYPE: DNA

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95	<213> ORGANISM: Arabidopsis thaliana	
96	<400> SEQUENCE: 5	
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99	<211> LENGTH: 29	
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101	<213> ORGANISM: Arabidopsis thaliana	
102	<400> SEQUENCE: 6	
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105	<211> LENGTH: 30	
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107	<213> ORGANISM: Arabidopsis thaliana	
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113	<213> ORGANISM: Arabidopsis thaliana	
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124	<212> TYPE: DNA	
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130	<212> TYPE: DNA	
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136	<212> TYPE: DNA	
137	<213> ORGANISM: Arabidopsis thaliana	
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140	<210> SEQ ID NO 13	
141	<211> LENGTH: 33	
142	<212> TYPE: DNA	
143	<213> ORGANISM: Arabidopsis thaliana	
144	<400> SEQUENCE: 13	

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159	<211> LENGTH: 28	
160	<212> TYPE: DNA	
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165	<211> LENGTH: 20	
166	<212> TYPE: DNA	
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171	<211> LENGTH: 26	
172	<212> TYPE: DNA	
173	<213> ORGANISM: Hordeum vulgare	
174	<400> SEQUENCE: 18	
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VERIFICATION SUMMARY
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Line ? Error/Warning

Original Text

19 W "N" or "Xaa" used: Feature required

gttgtaaaac gacggncagt gaattgtaat acgactcn

38 W "N" or "Xaa" used: Feature required

ttatcgatac cgtcgacctc gagggggggc ccggtacc